```
<!--StartFragment-->RESULT 2
AAD56890
ID
     AAD56890 standard; cDNA; 1279 BP.
XX
AC
    AAD56890;
XX
DT
     06-NOV-2003 (first entry)
XX
DE
     Human diacylqlycerol acyltransferase 2 (DGAT2) cDNA, 112023.
XX
KW
     Human; diacylglycerol acyltransferase 2; DGAT2; obesity; arrhythmia;
     coronary artery disease; hypertension; heart failure; tissue typing;
KW
     aberrant lipogenesis; cardiovascular disorder; atherosclerosis; angina;
ΚW
KW
     atrial fibrillation; dilated cardiomyopathy; idiopathic cardiomyopathy;
     diabetes; chromosome mapping; forensic biology; enzyme; gene; ss.
KW
XX
OS
     Homo sapiens.
XX
FH
     Key
                     Location/Qualifiers
FT
     CDS
                     42. .1028
FT
                     /*tag= a
FT
                     /product= "Human diacylqlycerol acyltransferase 2"
XX
PN
     WO2003053363-A2.
XX
PD
     03-JUL-2003.
XX
     19-DEC-2002; 2002WO-US040974.
PF
XX
PR
     19-DEC-2001; 2001US-0341947P.
PR
     19-SEP-2002; 2002US-0411859P.
XX
PΑ
     (MILL-) MILLENNIUM PHARM INC.
XX
ΡI
     Gimeno RE, Wu Z, Kapeller-Libermann R, Hubbard BK;
XX
DR
     WPI; 2003-559092/52.
DR
     P-PSDB; AAE37790.
XX
     New human diacylglycerol acyltransferase 2 (DGAT2) family member
PΤ
PΤ
     polypeptide and nucleic acid molecules, useful for diagnosing and
PT
     treating obesity, diabetes, atherosclerosis, aberrant lipogenesis or
PΤ
     triglyceride synthesis.
XX
PS
     Claim 1; Page 133-134; 154pp; English.
XX
CC
     The invention relates to human diacylglycerol acyltransferase 2 (DGAT2)
CC
     family members and their uses. DGAT2 family member sequences or their
CC
     modulators are useful for diagnosing and treating a subject with a
CC
     disorder associated with the aberrant DGAT family member polypeptide
     activity or nucleic acid expression, such as a disorder associated with
CC
CC
     obesity, diabetes, aberrant lipogenesis or triglyceride synthesis, or
CC
     cardiovascular disorder (e.g. atherosclerosis, coronary artery disease,
CC
     hypertension, heart failure, atrial fibrillation, arrhythmias, dilated
CC
     cardiomyopathy, idiopathic cardiomyopathy or angina). The invention is
CC
     also useful in screening assays (e.g. tissue typing, chromosome mapping,
CC
     or in forensic biology), in predictive medicine (e.g. diagnostic assays,
CC
     prognostic assays, monitoring clinical trials or pharmacogenetics), or as
CC
     surrogate markers (e.g. markers of disease states or markers of drug
CC
     activity). The present sequence is human DGAT2 cDNA
XX
```

```
Sequence 1279 BP; 273 A; 352 C; 328 G; 326 T; 0 U; 0 Other;
                       Score 1084.8;
                  95.9%;
 Query Match
                                DB 2;
                                     Length 1279;
 Best Local Similarity
                  98.0%;
 Matches 1109; Conservative
                      0;
                        Mismatches
                                  22;
                                      Indels
                                             1;
                                               Gaps
                                                     1;
        1 ACTGTTCTGAGATCTTTGCCTCCCTCAGGCTCCCGAGAATCATGGCTCATTCCAAGCAGC 60
Qу
         1 ACTGTTCTGAGATCTTTGCCTCCCTCAGGCTCCCGAGAATCATGGCTCATTCCAAGCAGC 60
Db
       61 CTAGTCACTTCCAGAGTCTGATGCTTCTGCAGTGGCCTTTGAGCTACCTTGCCATCTTTT 120
Qу
         61 CTAGTCACTTCCAGAGTCTGATGCTTCTGCAGTGGCCTTTGAGCTACCTTGCCATCTTTT 120
Db
      121 GGATCTTGCAGCCATTGTTCGTCTACCTGCTGTTTACATCCTTGTGGCCGCTACCAGTGC 180
Qу
         121 GGATCTTGCAGCCATTGTTCGTCTACCTGCTGTTTACATCCTTGTGGCCGCTACCAGTGC 180
Db
      181 TTTACTTTGCCTGGTTGTTCCTGGACTGGAAGACCCCAGAGCGAGGTGGCAGGCGTTCGG 240
Qу
         181 TTTACTTTGCCTGGTTGTTCCTGGACTGGAAGACCCCAGAGCGAGGTGGCAGGCGTTCGG 240
Db
       241 CCTGGGTAAGGAACTGGTGTGTCTGGACCCACATCAGGGACTATTTCCCCATTACGATCC 300
Qу
         241 CCTGGGTAAGGAACTGGTGTGTCTGGACCCACATCAGGGACTATTTCCCCATTACGATCC 300
Db
      301 TGAAGACAAAGGACCTATCACCTGAGCACAACTACCTCATGGGGGTTCACCCCCATGGCC 360
Qу
         301 TGAAGACAAAGGACCTATCACCTGAGCACAACTACCTCATGGGGGTTCACCCCCATGGCC 360
Db
       361 TCCTGACCTTTGGCGCCTTCTGCAACTTCTGCACTGAGGCCACAGGCTTCTCGAAGACCT 420
Qу
         361 TCCTGACCTTTGGCGCCTTCTGCAACTTCTGCACTGAGGCCACAGGCTTCTCGAAGACCT 420
Db
       421 TCCCAGGCATCACTCCTCACTTGGCCACGCTGTCCTGGTTCTTCAAGATCCCCTTTGTTA 480
Qу
         421 TCCCAGGCATCACTCCTCACTTGGCCACACTGTCCTGGTTCTTCAAGATCCCCTTTGTTA 480
Db
       Qу
         Db
Qу
      541 TGAGCCATGGCACTGGCAACCTCGTGGGCATTGTAGTGGGAGGTGTGGGTGAGGCCCTGC 600
         541 TGAGCCATGGCACTGGCAACCTCGTGGGCATTGTAGTGGGAGGTGTGGGTGAGGCCCTGC 600
Db
       601 AAAGTGTGCCCAACACCACCACCTCATCCTCCAGAAGCGCAAGGGGTTCGTGCGCACAG 660
Qу
         601 AAAGTGTGCCCAAGACCACCACCCTCATCCTCCAGAAGCGCAAGGGGTTCGTGCGCACAG 660
Db
       661 CCCTCCAGCATGGGGCTCATCTGGTCCCCACCTTCACTTTTGGGGAAACTGAGGTGTATG 720
Qу
         661 CCCTCCAGCATGGGGCTCATCTGGTCCCCACCTTCACTTTTGGGGAAACTGAGGTGTATG 720
Db
      721 ATCAGGTGCTGTTCCATAAGGATAGCAGGATGTACAAGTTCCAGAGCTGCTTCCGCCGTA 780
Qу
         721 ATCAGGTGCTGTTCCATAAGGATAGCAGGATGTACAAGTTCCAGAGCTGCTTCCGCCGTA 780
Db
      781 TCTTTGGTTTCTACTGTTGTGTCTTCTATGGACAAGCTTCTGTCAAGGCTCCACTGGGC 840
Qу
         781 TCTTTGGTTTCTACTGTTGTGTCTTCTATGGACAAAGCTTCTGTCAAGGCTCCACTGGGC 840
Db
```

Qу	841	TCCTGCCATACTCCAGGCCTATTGTCACTGTGGTTGGGGAGCCTCTGCCACTGCCCCAAA	900
Db	841	TCCTGCCATACTCCAGGCCTATTGTCACTGTGGTTGGGGAGCCTCTGCCACTGCCCCAAA	900
Qy	901	TTGAAAAGCCAAGCCAGGAGATGGTGGACAAATACCATGCACTTTATATGGATGCTCTGC	960
Db	901	TTGAAAAGCCAAGCCAGGAGATGGTGGACAAATACCATGCACTTTATATGGATGCTCTGG	960
Qy	961	ACAAACTGTTCGACCAGCATAAGACCCACTATGGCTGCTCAGAGACCCAAAAGCTGTTTT	1020
Db	961	ACAAACTGTTCGACCAGCATAAGACCCACTATGGCTGCTCAGAGACCCAAAAGCTGTTTT	1020
Qу	1021	TCCTGTGAATGAAGGTACTGCATGCCCAGGAGCACAGGAGTGCCTGCC	1079
Db	1021	TCCTGTGAATGAAGGTACTGCATGCCCAGGAGCACAGGAGTGCCTGCC	1080
QУ	1080	ACTCATCTGCCACTAACCAAAGACAGGCAGGAGATGAGGGAGG	
Db 1081 GAATCATCTGGCATAACCAAAGACAGGCAGGAGATGGAGGGAG			